

+	1	MENGTHVNGSANGFCIKDPLNWGVAAEALTGSHLDEVKKMVGEFRKPVVK	Sunflower
	1	MENGTHVNGVNELCIKDPLNWGVAAEALTGSHLDEVKKMVAEFRKPVVK	100%
	51	LGGETLTVSQVAGISAAGDGNMVKVELSEARAGVKASSDWVME SMNKGT	Sunflower
	51	LGGETLTVSQVAGIAAANDSDTVKVELSEARAGVKASSDWVME SMNKGT	100%
	101	DSYGVTTGFGATSHRRTKNGGALQKELIRFLNAGIFGNGTESSHTLPHSA	Sunflower
	101	DSYGVTTGFGATSHRRTKGGALQKELIRFLNAGIFGNGTEHTLPHSA	100%
	151	TRAAMIVRINTLLQGYSGIRFEILEAITKFLNNNITPCLPLRGTTITASGD	Sunflower
	151	TRAAMIVRINTLLQGYSGIRFEILEAITKFLN N TP LPLRGTTITASGD	100%
	201	LVPLSYIAGLLTGRPNASKAVGPAGEVLNAESAFAAGVEGGFFELQPKEG	Sunflower
	201	LVPLSYIAGLLTGRNSKAVGPTGEVLNAEKAFAGVEGGFFELQPKEG	100%
	251	LALVNGTAVGSGMASMVLFEANVLALLSEVLSAIFAEMVQCKPEFTDHLT	Sunflower
	251	LALVNGTAVGSGMASMVLFDANVLALLSEVLSAIFAEMVQCKPEFTDHLT	100%
	301	HKLKHHHPGQIEAAAIMHEYILDGSDYVKAQKVHEMDPLQKPKQDRYALRT	Sunflower
	301	HKLKHHHPGQIEAAAIMHEYILDGSDYVKAQKVHEMDPLQKPKQDRYALRT	100%
	351	SPQWLGPQIEVIRSATKMIEREINSVNDNPLIDVSRNKALHGGNFQGTPI	Sunflower
	351	SPQWLGPQIEVIRSSTKMIEREINSVNDNPLIDVSRNKALHGGNFQGTPI	100%
	401	GVSMNTRLAIAAIGKVTIAQFSELVNDFYNNGLPSHLSGGRNPSLDSCGF	Sunflower
	401	GVSMNTRLAIAAIGKLMFAQFSELVNDFYNNGLPSNLSGGRNPSLDYGF	100%
	451	KGGEIAMASYCSELQFLANPVTNHVQSAEQHNQDVNSLGLISARKTAEAV	Sunflower
	451	KGGEIAMASYCSELQFLANPVTNHVQSAEQHNQDVNSLGLISARKTAEAV	100%
	501	DILKLMSSSTYLVALCQSIDLRHLEENMKSTVKNTVSQVAKKVLTMGVNGE	Sunflower
	501	DILKLMSSSTYLVALCQSIDLRHLEENMKSTVKNTVSQVAKKVLTMGVNGE	100%
	551	LHPSRFCEKDLLRVVDREYVFAYADDPCLTTYPLMQKLRQVLVDHALNNG	Sunflower
	551	LHPSRFCEKDLLRVVDREYVFAYIDDVCSGTYPLMQKLRQVLVDHALNNG	100%
	601	ETEKNTSIFQKIATFEDELKAILLPKEVESVRVAFENGTM SIPNRIKAC	Sunflower
	601	ETEKNTSIFQKIATFEDELKVLLPKEVEGVRVAYENDT SIPNRIKAC	100%
	651	RSYPLYRFVREELGGA	Sunflower
	651	RSYPLYRFVREELGRGFLTGEKVTSPGEEFDRVFTAMCKGQIIDPLECL	100%
	701	GGWNGEPLPIC	100%

Figure 3